

SEQUENCE LISTING

<110> Chiquet-Ehrismann, Ruth
Scherberich, Arnaud

<120> TENASCIN-W COMPOSITIONS AND USES THEREOF

<130> 1/32411A/USN/FMI

<140> 10/509,009

<141> 2004-09-24

<150> PCT/EP03/03150

<151> 2003-03-26

<150> GB0207224.7

<151> 2002-03-27

<160> 28

<170> FastSEQ for Windows Version 4.0

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<212> DNA

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<221> CDS

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<223> Mouse tenascin-W

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Asp Gly Glu Thr Lys Glu Lys Ala Val Pro Lys Asp Gln Ser Ser Thr	
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Pro	Asp	Cys	Ser	Gln	Val	Val	Ala	Pro	Gln	Gly	Leu	Gln	Leu	Leu	Lys
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Leu	Leu	Pro	Gly	Thr	Lys	Tyr	Ile	Val	Thr	Leu	Arg	Asn	Val	Lys	Lys
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Thr Glu Ile Asp Ser Pro	Glu Asn Leu Val Thr Asp Arg Val Thr Glu			
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Val Glu Tyr Lys Val Tyr	Val Trp Ala Glu Lys Gly Asp Arg Glu Ser			
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Lys Lys Ala Asn Thr Lys	Ala Pro Thr Asp Ile Asp Ser Pro Lys Asn			
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Leu Val Thr Asp Gln Val	Thr Glu Asn Thr Leu Ser Val Ser Trp Asp			
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Pro Val Gln Ala Asn Ile	Asp Arg Tyr Met Val Ser Tyr Thr Ser Ala			
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Asp Gly Glu Thr Arg Glu	Val Pro Val Pro Lys Glu Lys Ser Ser Thr			
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Val Leu Thr Gly Leu Arg	Pro Gly Val Glu Tyr Lys Val His Val Trp			
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Ala Gln Lys Gly Thr Gln	Glu Ser Arg Lys Ala Asn Thr Lys Ala Pro			
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Thr Asp Ile Asp Gly Pro	Lys Asn Leu Val Thr Asp Gln Val Thr Glu			
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Thr Thr Leu Ser Val Ser	Trp Asp Pro Val Glu Ala Asp Ile Asp Arg			
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Tyr Met Val Arg Tyr Thr	Ser Pro Asp Gly Glu Thr Lys Glu Val Pro			
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Val Ser Lys Asp Lys Ser	Ser Thr Val Leu Arg Gly Leu Arg Pro Gly			
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Val Glu Tyr Lys Val Asp	Val Trp Ala Gln Lys Gly Ala Gln Asp Ser			
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Arg Lys Ala Asn Thr Lys	Ala Pro Thr Asp Ile Asp Ser Pro Lys Asn			
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Pro Val Glu Ala Asp Ile	Asp Arg Tyr Val Val Arg Tyr Thr Ser Ala			
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Asp Gly Glu Thr Arg Glu	Ile Pro Val Arg Lys Glu Lys Ser Ser Thr			
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Val Leu Thr Gly Leu Arg	Pro Gly Val Glu Tyr Thr Val Gln Val Trp			
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Ala Gln Lys Gly Ala Arg	Glu Ser Lys Lys Ala Lys Thr Lys Ala Pro			
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Thr Glu Ile Asp Ser Pro	Lys Asn Leu Val Thr Asn Arg Val Thr Glu			
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Asn Thr Ala Thr Ile Ser	Trp Asp Pro Val Arg Ala Asn Ile Asp Arg			
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Tyr Met Val Arg Tyr Thr	Ser Ala Asp Gly Glu Thr Lys Glu Ile Pro			
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Phe	Arg	His	Asn	Ile	Arg	Leu	Gln	Thr	Pro	Gln	Lys	Asp	Cys	Glu	Leu	
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Lys	Pro	Gly	Glu	Ala	Tyr	Lys	Val	Tyr	Val	Trp	Ala	Glu	Arg	Gly	Asn	

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cgc	tac	acc	tct	gct	gac	gga	gag	acc	agg	gag	gtt	ccg	gtg	ggg	aag	2784
Arg	Tyr	Thr	Ser	Ala	Asp	Gly	Glu	Thr	Arg	Glu	Val	Pro	Val	Gly	Lys	
		915					920					925				
gag	cac	agc	agc	act	gtc	ctg	acg	ggc	ctg	aga	cca	ggc	atg	gag	tac	2832
Glu	His	Ser	Ser	Thr	Val	Leu	Thr	Gly	Leu	Arg	Pro	Gly	Met	Glu	Tyr	
	930					935					940					
atg	gtg	cac	gtg	tgg	gcc	cag	aag	ggg	gcc	cag	gag	agc	aag	aag	gct	2880
Met	Val	His	Val	Trp	Ala	Gln	Lys	Gly	Ala	Gln	Glu	Ser	Lys	Lys	Ala	
	945				950					955					960	
gac	acc	aag	gcc	cag	aca	gaa	ctc	gac	cct	ccc	aga	aac	ctt	cgt	cca	2928
Asp	Thr	Lys	Ala	Gln	Thr	Glu	Leu	Asp	Pro	Pro	Arg	Asn	Leu	Arg	Pro	
				965					970					975		
tct	gct	gta	acg	cag	tct	ggt	ggc	ata	ttg	acc	tgg	acg	ccc	ccc	tct	2976
Ser	Ala	Val	Thr	Gln	Ser	Gly	Gly	Ile	Leu	Thr	Trp	Thr	Pro	Pro	Ser	
			980					985					990			

gct cag atc cac ggc tac att ctg act tac cag ttc cca gat ggc aca Ala Gln Ile His Gly Tyr Ile Leu Thr Tyr Gln Phe Pro Asp Gly Thr 995 1000 1005	3024
gtt aag gag atg cag ctg gga cgg gaa gac cag agg ttt gcg ttg caa Val Lys Glu Met Gln Leu Gly Arg Glu Asp Gln Arg Phe Ala Leu Gln 1010 1015 1020	3072
ggc ctt gag caa ggc gcc acc tac cct gtc tcc ctt gtt gcc ttt aag Gly Leu Glu Gln Gly Ala Thr Tyr Pro Val Ser Leu Val Ala Phe Lys 1025 1030 1035 1040	3120
ggg ggt cgc cgg agc aga aat gta tcc acc acc ctc tcc aca gtt ggt Gly Gly Arg Arg Ser Arg Asn Val Ser Thr Thr Leu Ser Thr Val Gly 1045 1050 1055	3168
gcc cgt ttc cca cac cct tcg gac tgc agt cag gtt cag cag aac agc Ala Arg Phe Pro His Pro Ser Asp Cys Ser Gln Val Gln Gln Asn Ser 1060 1065 1070	3216
aat gcc gcc agt ggt ctg tac acc atc tac ctg cat ggc gat gcc agc Asn Ala Ala Ser Gly Leu Tyr Thr Ile Tyr Leu His Gly Asp Ala Ser 1075 1080 1085	3264
cgg ccc ctg cag gtg tac tgt gac atg gaa acg gac gga ggt ggc tgg Arg Pro Leu Gln Val Tyr Cys Asp Met Glu Thr Asp Gly Gly Gly Trp 1090 1095 1100	3312
att gtc ttc cag agg cgg aac act ggg cag ctg gat ttc ttc aag cga Ile Val Phe Gln Arg Arg Asn Thr Gly Gln Leu Asp Phe Phe Lys Arg 1105 1110 1115 1120	3360
tgg agg agc tat gtg gaa ggc ttt ggg gac ccc atg aag gag ttc tgg Trp Arg Ser Tyr Val Glu Gly Phe Gly Asp Pro Met Lys Glu Phe Trp 1125 1130 1135	3408
ctt gga ctt gac aag cta cac aac ctc acc acc ggc act cca gcg cgg Leu Gly Leu Asp Lys Leu His Asn Leu Thr Thr Gly Thr Pro Ala Arg 1140 1145 1150	3456
tat gag gtg aga gtg gat tta cag act gcc aat gaa tct gcc tat gct Tyr Glu Val Arg Val Asp Leu Gln Thr Ala Asn Glu Ser Ala Tyr Ala 1155 1160 1165	3504
ata tat gat ttc ttc caa gtg gcc tcc agc aag gag cgg tat aag ctg Ile Tyr Asp Phe Phe Gln Val Ala Ser Ser Lys Glu Arg Tyr Lys Leu 1170 1175 1180	3552
aca gtt ggg aaa tac aga ggc acg gca ggg gat gct ctt act tac cac Thr Val Gly Lys Tyr Arg Gly Thr Ala Gly Asp Ala Leu Thr Tyr His 1185 1190 1195 1200	3600
aat gga tgg aag ttt aca act ttt gac aga gac aat gat atc gca ctc Asn Gly Trp Lys Phe Thr Thr Phe Asp Arg Asp Asn Asp Ile Ala Leu 1205 1210 1215	3648
agc aac tgt gcc ctg aca cat cat ggt ggc tgg tgg tat aag aac tgc Ser Asn Cys Ala Leu Thr His His Gly Gly Trp Trp Tyr Lys Asn Cys 1220 1225 1230	3696
cac ttg gcc aac cct aat ggc aga tat ggg gag acc aag cac agt gag	3744

His	Leu	Ala	Asn	Pro	Asn	Gly	Arg	Tyr	Gly	Glu	Thr	Lys	His	Ser	Glu		
		1235					1240					1245					
ggg	gtg	aac	tgg	gag	cct	tgg	aaa	gga	cat	gaa	ttc	tcc	att	cct	tac	3792	
Gly	Val	Asn	Trp	Glu	Pro	Trp	Lys	Gly	His	Glu	Phe	Ser	Ile	Pro	Tyr		
		1250				1255					1260						
gtg	gag	ttg	aaa	atc	cgc	cct	cat	ggc	tac	agc	agg	gag	cct	gtc	ctg	3840	
Val	Glu	Leu	Lys	Ile	Arg	Pro	His	Gly	Tyr	Ser	Arg	Glu	Pro	Val	Leu		
		1265			1270					1275					1280		
ggc	aga	aag	aag	cgg	acg	ctg	aga	gga	agg	ctg	cga	acg	ttc	tga		3885	
Gly	Arg	Lys	Lys	Arg	Thr	Leu	Arg	Gly	Arg	Leu	Arg	Thr	Phe	*			
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 <213> Homo sapiens

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Ala	Ser	Ala	Pro	Ala	Thr	Leu	Glu	Pro	Pro	Gly	Cys	Ser	Asn	Lys	Glu	
		20						25					30			
Gln	Gln	Val	Thr	Val	Ser	His	Thr	Tyr	Lys	Ile	Asp	Val	Pro	Lys	Ser	
		35					40					45				
Ala	Leu	Val	Gln	Val	Asp	Ala	Asp	Pro	Gln	Pro	Leu	Ser	Asp	Asp	Gly	
	50					55				60						
Ala	Ser	Leu	Leu	Ala	Leu	Gly	Glu	Ala	Arg	Glu	Glu	Gln	Asn	Ile	Ile	
65					70				75					80		
Phe	Arg	His	Asn	Ile	Arg	Leu	Gln	Thr	Pro	Gln	Lys	Asp	Cys	Glu	Leu	
			85					90					95			
Ala	Gly	Ser	Val	Gln	Asp	Leu	Leu	Ala	Arg	Val	Lys	Lys	Leu	Glu	Glu	
			100					105					110			
Glu	Met	Val	Glu	Met	Lys	Glu	Gln	Cys	Ser	Ala	Gln	Arg	Cys	Cys	Gln	
		115					120					125				
Gly	Val	Thr	Asp	Leu	Ser	Arg	His	Cys	Ser	Gly	His	Gly	Thr	Phe	Ser	
	130					135					140					
Leu	Glu	Thr	Cys	Ser	Cys	His	Cys	Glu	Glu	Gly	Arg	Glu	Gly	Pro	Ala	
145					150					155					160	
Cys	Glu	Arg	Leu	Ala	Cys	Pro	Gly	Ala	Cys	Ser	Gly	His	Gly	Arg	Cys	
			165					170						175		
Val	Asp	Gly	Arg	Cys	Leu	Cys	His	Glu	Pro	Tyr	Val	Gly	Ala	Asp	Cys	
			180					185					190			
Gly	Tyr	Pro	Ala	Cys	Pro	Glu	Asn	Cys	Ser	Gly	His	Gly	Glu	Cys	Val	
		195					200						205			
Arg	Gly	Val	Cys	Gln	Cys	His	Glu	Asp	Phe	Met	Ser	Glu	Asp	Cys	Ser	
		210				215					220					
Glu	Lys	Arg	Cys	Pro	Gly	Asp	Cys	Ser	Gly	His	Gly	Phe	Cys	Asp	Thr	
225					230					235					240	
Gly	Glu	Cys	Tyr	Cys	Glu	Glu	Gly	Phe	Thr	Gly	Leu	Asp	Cys	Ala	Gln	
				245					250					255		
Val	Val	Thr	Pro	Gln	Gly	Leu	Gln	Leu	Leu	Lys	Asn	Thr	Glu	Asp	Ser	
			260					265					270			
Leu	Leu	Val	Ser	Trp	Glu	Pro	Ser	Ser	Gln	Val	Asp	His	Tyr	Leu	Leu	
		275					280					285				
Ser	Tyr	Tyr	Pro	Leu	Gly	Lys	Glu	Leu	Ser	Gly	Lys	Gln	Ile	Gln	Val	
		290				295					300					

Pro	Lys	Glu	Gln	His	Ser	Tyr	Glu	Ile	Leu	Gly	Leu	Leu	Pro	Gly	Thr	305	310	315	320
Lys	Tyr	Ile	Val	Thr	Leu	Arg	Asn	Val	Lys	Asn	Glu	Val	Ser	Ser	Ser	325	330	335	
Pro	Gln	His	Leu	Leu	Ala	Thr	Thr	Asp	Leu	Ala	Val	Leu	Gly	Thr	Ala	340	345	350	
Trp	Val	Thr	Asp	Glu	Thr	Glu	Asn	Ser	Leu	Asp	Val	Glu	Trp	Glu	Asn	355	360	365	
Pro	Ser	Thr	Glu	Val	Asp	Tyr	Tyr	Lys	Leu	Arg	Tyr	Gly	Pro	Met	Thr	370	375	380	
Gly	Gln	Glu	Val	Ala	Glu	Val	Thr	Val	Pro	Lys	Ser	Ser	Asp	Pro	Lys	385	390	395	400
Ser	Arg	Tyr	Asp	Ile	Thr	Gly	Leu	His	Pro	Gly	Thr	Glu	Tyr	Lys	Ile	405	410	415	
Thr	Val	Val	Pro	Met	Arg	Gly	Glu	Leu	Glu	Gly	Lys	Pro	Ile	Leu	Leu	420	425	430	
Asn	Gly	Arg	Thr	Glu	Ile	Asp	Ser	Pro	Thr	Asn	Val	Val	Thr	Asp	Arg	435	440	445	
Val	Thr	Glu	Asp	Thr	Ala	Thr	Val	Ser	Trp	Asp	Pro	Val	Gln	Ala	Val	450	455	460	
Ile	Asp	Lys	Tyr	Val	Val	Arg	Tyr	Thr	Ser	Ala	Asp	Gly	Asp	Thr	Lys	465	470	475	480
Glu	Met	Ala	Val	His	Lys	Asp	Glu	Ser	Ser	Thr	Val	Leu	Thr	Gly	Leu	485	490	495	
Lys	Pro	Gly	Glu	Ala	Tyr	Lys	Val	Tyr	Val	Trp	Ala	Glu	Arg	Gly	Asn	500	505	510	
Gln	Gly	Ser	Lys	Lys	Ala	Asp	Thr	Asn	Ala	Leu	Thr	Glu	Ile	Asp	Ser	515	520	525	
Pro	Ala	Asn	Leu	Val	Thr	Asp	Arg	Val	Thr	Glu	Asn	Thr	Ala	Thr	Ile	530	535	540	
Ser	Trp	Asp	Pro	Val	Gln	Ala	Thr	Ile	Asp	Lys	Tyr	Val	Val	Arg	Tyr	545	550	555	560
Thr	Ser	Ala	Asp	Asp	Gln	Glu	Thr	Arg	Glu	Val	Leu	Val	Gly	Lys	Glu	565	570	575	
Gln	Ser	Ser	Thr	Val	Leu	Thr	Gly	Leu	Arg	Pro	Gly	Val	Glu	Tyr	Thr	580	585	590	
Val	His	Val	Trp	Ala	Gln	Lys	Gly	Asp	Arg	Glu	Ser	Lys	Lys	Ala	Asp	595	600	605	
Thr	Asn	Ala	Pro	Thr	Asp	Ile	Asp	Ser	Pro	Lys	Asn	Leu	Val	Thr	Asp	610	615	620	
Arg	Val	Thr	Glu	Asn	Met	Ala	Thr	Val	Ser	Trp	Asp	Pro	Val	Gln	Ala	625	630	635	640
Ala	Ile	Asp	Lys	Tyr	Val	Val	Arg	Tyr	Thr	Ser	Ala	Gly	Gly	Glu	Thr	645	650	655	
Arg	Glu	Val	Pro	Val	Gly	Lys	Glu	Gln	Ser	Ser	Thr	Val	Leu	Thr	Gly	660	665	670	
Leu	Arg	Pro	Gly	Met	Glu	Tyr	Met	Val	His	Val	Trp	Ala	Gln	Lys	Gly	675	680	685	
Asp	Gln	Glu	Ser	Lys	Lys	Ala	Asp	Thr	Lys	Ala	Gln	Thr	Asp	Ile	Asp	690	695	700	
Ser	Pro	Gln	Asn	Leu	Val	Thr	Asp	Arg	Val	Thr	Glu	Asn	Met	Ala	Thr	705	710	715	720
Val	Ser	Trp	Asp	Pro	Val	Arg	Ala	Thr	Ile	Asp	Arg	Tyr	Val	Val	Arg	725	730	735	
Tyr	Thr	Ser	Ala	Lys	Asp	Gly	Glu	Thr	Arg	Glu	Val	Pro	Val	Gly	Lys	740	745	750	
Glu	Gln	Ser	Ser	Thr	Val	Leu	Thr	Gly	Leu	Arg	Pro	Gly	Val	Glu	Tyr	755	760	765	
Thr	Val	His	Val	Trp	Ala	Gln	Lys	Gly	Ala	Gln	Glu	Ser	Lys	Lys	Ala	770	775	780	
Asp	Thr	Lys	Ala	Gln	Thr	Asp	Ile	Asp	Ser	Pro	Gln	Asn	Leu	Val	Thr				

785					790					795					800
Asp	Trp	Val	Thr	Glu	Asn	Thr	Ala	Thr	Val	Ser	Trp	Asp	Pro	Val	Gln
				805					810					815	
Ala	Thr	Ile	Asp	Arg	Tyr	Val	Val	His	Tyr	Thr	Ser	Ala	Asn	Gly	Glu
			820					825					830		
Thr	Arg	Glu	Val	Pro	Val	Gly	Lys	Glu	Gln	Ser	Ser	Thr	Val	Leu	Thr
			835				840					845			
Gly	Leu	Arg	Pro	Gly	Met	Glu	Tyr	Thr	Val	His	Val	Trp	Ala	Gln	Lys
	850					855					860				
Gly	Asn	Gln	Glu	Ser	Lys	Lys	Ala	Asp	Thr	Lys	Ala	Gln	Thr	Glu	Ile
865					870					875				880	
Asp	Gly	Pro	Lys	Asn	Leu	Val	Thr	Asp	Trp	Val	Thr	Glu	Asn	Met	Ala
				885					890					895	
Thr	Val	Ser	Trp	Asp	Pro	Val	Gln	Ala	Thr	Ile	Asp	Lys	Tyr	Met	Val
			900					905					910		
Arg	Tyr	Thr	Ser	Ala	Asp	Gly	Glu	Thr	Arg	Glu	Val	Pro	Val	Gly	Lys
			915				920					925			
Glu	His	Ser	Ser	Thr	Val	Leu	Thr	Gly	Leu	Arg	Pro	Gly	Met	Glu	Tyr
	930					935					940				
Met	Val	His	Val	Trp	Ala	Gln	Lys	Gly	Ala	Gln	Glu	Ser	Lys	Lys	Ala
945					950					955					960
Asp	Thr	Lys	Ala	Gln	Thr	Glu	Leu	Asp	Pro	Pro	Arg	Asn	Leu	Arg	Pro
			965						970					975	
Ser	Ala	Val	Thr	Gln	Ser	Gly	Gly	Ile	Leu	Thr	Trp	Thr	Pro	Pro	Ser
			980					985					990		
Ala	Gln	Ile	His	Gly	Tyr	Ile	Leu	Thr	Tyr	Gln	Phe	Pro	Asp	Gly	Thr
			995				1000					1005			
Val	Lys	Glu	Met	Gln	Leu	Gly	Arg	Glu	Asp	Gln	Arg	Phe	Ala	Leu	Gln
	1010					1015					1020				
Gly	Leu	Glu	Gln	Gly	Ala	Thr	Tyr	Pro	Val	Ser	Leu	Val	Ala	Phe	Lys
1025					1030					1035					1040
Gly	Gly	Arg	Arg	Ser	Arg	Asn	Val	Ser	Thr	Thr	Leu	Ser	Thr	Val	Gly
				1045					1050					1055	
Ala	Arg	Phe	Pro	His	Pro	Ser	Asp	Cys	Ser	Gln	Val	Gln	Gln	Asn	Ser
			1060					1065					1070		
Asn	Ala	Ala	Ser	Gly	Leu	Tyr	Thr	Ile	Tyr	Leu	His	Gly	Asp	Ala	Ser
			1075				1080					1085			
Arg	Pro	Leu	Gln	Val	Tyr	Cys	Asp	Met	Glu	Thr	Asp	Gly	Gly	Gly	Trp
	1090					1095					1100				
Ile	Val	Phe	Gln	Arg	Arg	Asn	Thr	Gly	Gln	Leu	Asp	Phe	Phe	Lys	Arg
1105					1110					1115					1120
Trp	Arg	Ser	Tyr	Val	Glu	Gly	Phe	Gly	Asp	Pro	Met	Lys	Glu	Phe	Trp
				1125					1130					1135	
Leu	Gly	Leu	Asp	Lys	Leu	His	Asn	Leu	Thr	Thr	Gly	Thr	Pro	Ala	Arg
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Tyr	Glu	Val	Arg	Val	Asp	Leu	Gln	Thr	Ala	Asn	Glu	Ser	Ala	Tyr	Ala
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Ile	Tyr	Asp	Phe	Phe	Gln	Val	Ala	Ser	Ser	Lys	Glu	Arg	Tyr	Lys	Leu
	1170					1175					1180				
Thr	Val	Gly	Lys	Tyr	Arg	Gly	Thr	Ala	Gly	Asp	Ala	Leu	Thr	Tyr	His
1185					1190					1195					1200
Asn	Gly	Trp	Lys	Phe	Thr	Thr	Phe	Asp	Arg	Asp	Asn	Asp	Ile	Ala	Leu
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Ser	Asn	Cys	Ala	Leu	Thr	His	His	Gly	Gly	Trp	Trp	Tyr	Lys	Asn	Cys
			1220					1225					1230		
His	Leu	Ala	Asn	Pro	Asn	Gly	Arg	Tyr	Gly	Glu	Thr	Lys	His	Ser	Glu
			1235				1240					1245			
Gly	Val	Asn	Trp	Glu	Pro	Trp	Lys	Gly	His	Glu	Phe	Ser	Ile	Pro	Tyr
	1250					1255					1260				
Val	Glu	Leu	Lys	Ile	Arg	Pro	His	Gly	Tyr	Ser	Arg	Glu	Pro	Val	Leu
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Gly Arg Lys Lys Arg Thr Leu Arg Gly Arg Leu Arg Thr Phe
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<220>
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<210> 6
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<223> First PCR primer for sequencing 5' end

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<210> 10
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<223> First PCR reaction for sequencing 3' end

<400> 13
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<210> 15
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<400> 15
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<210> 16
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<210> 17
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<213> Homo sapiens

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<220>
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29